

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 116201

TO: Nita M Minnifield Location: REM-3C18

Art Unit: 1645

Monday, March 08, 2004

Case Serial Number: 09/056019

From: Barb O'Bryen

Location: Biotech-Chem Library

Remsen E01A69

Phone: 571-272-2518

SOR

barbara.obryen@uspto.gov

Search Notes	
}	



STIC-Biotech/ChemLib

116201

From:

Chan, Christina

Sent:

Monday, March 08, 2004 8:37 AM

To: Subject: Minnifield, Nita; Pak, Michael; STIC-Biotech/ChemLib

RE: rush sequence search request

Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89

----Original Message----

From:

Minnifield, Nita.

Sent:

Sunday, March 07, 2004 10:43 AM

To:

Chan, Christina

Subject:

rush sequence search request

Christina, please approve, 2 month overdue amdt.

STIC

09/056019

Please do commercial and interference sequence search on SEQ ID NO: 1, 3-5, 7, 9-11, 22-24 (all amino acid sequences).

Please provide a paper copy of results.

Transferred case

Thanks, Nita M. Minnifield

Searcher:
Phone:
_ocation:
Date Picked Up:
Date Completed:
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST (where applic.)
STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.:
WWW/Internet:
Other (specify):

Art Unit 1645 Office REM-3C01 Mailbox REM-3C18 571-272-0860

Searcher:	
Phone:	_
Location:	
Date Picked Up:	
Date Completed:	
Searcher Prep/Review:	
Clerical:	_
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TYPE OF SEARCH:	
NA Sequences:	
AA Sequences:	
Structures:	
Bibliographic:	
Litigation:	
Full text:	
Patent Family:	
Other:	

VENDOR/COST (where applic.)
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Questel/Orbit:
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Lexis/Nexis:
Sequence Sys.:
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Other (specify):

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to the EMBL/GenBank/DDBJ databases.
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InterPro: IPR002479, CW binding.
InterPro: IPR005877, Gpos YSIRK.
InterPro: IPR007756, RICH.
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                                                                                                                                                                                                                                                                                                Tannelli F., Oggioni M.R., Pozzi G.,
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Bacteria, Firmicutes, Lactobacillales, Streptococcaceae,
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"Allelic Variation in the Highly Polymorphic Locus pspC
Streptococcus pneumoniae.";
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          Last annotation update)
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Pfam; PF05062; RICH; 2.
Pfam; PF05062; RICH; 2.
TIGRFAMS; TIGRO1167; LiPXTG anchor; 1.
TIGRFAMS; TIGRO1168; YSTRK signal; 1.
PROSITE; PS50847; GRAM POS ANCHORING; 1.
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04; Conservative 2
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   01-OCT-2003 (TrEMBLrel. 25,
                                                                                                       Streptococcus pneumoniae
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NCBI_TaxID=1313;
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96 KKTVDIVNKLONINNEYLNKIIQSTSTYBELQKLAMBSQSBVDKAVSEFEKDLSSSSSG 155
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                                                                                                                                                                                                                                                                                                                  Gaps
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STRAIN=serotype 1;
MEDLINE=98010350; PubMed=9350867;
Hammerschmidt S., Talay S.R., Brandtzaeg P., Chhatwal G.S.;
"SpsA, a novel pneumococcal surface protein with specific binding to Immunoglobulin A and secretory component.";
MMO1. Microbiol. 25:1113-1124 (1997).
EMBL; Y10818; CAA71783.1;
                     TO THE CELL WALL PEPTIDOGLYCAN BY
                                                                                                                                                                                                                                                                                                                  10;
                                                                                                                                                                                                                                                                                  Length 769;
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Local Similarity 73.6%; Pred. No. 7.8e-39;
les 203; Conservative 25; Mismatches 38; Indels
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Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
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Last annotation update)
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Pfam; PF01473; CW binding_1; 9.
Pfam; PF05062; RICH; 1.
Pfam; PF04650; YSIRK signal; 1.
TIGRFAMS; TIGR01168; YSIRK signal; 1.
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us-09-056-019b-3.rspt

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Surface protein PSpC.
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NCBI_TaxID=1313;
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                                                                                                                                         Gaps
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                                                                                                   Length 523;
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Barteria, Firmicutes; Lactobacillales; Streptococcaceae; Streptococcus, NCBI_TaxID=1313;
                  1 37 POTENTIAL.
38 523 POTENTIAL.
523 AA; 59151 MW; B3390DF8A706465A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Created)
Last sequence update)
Last annotation update)
                                                                                             Query Match 59.8%; Score 836; DB 2; Best Local Similarity 62.1%; Pred. No. 2.1e-33; Matches 182; Conservative 39; Mismatches 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-0CT-2000 (TrEMBLrel. 15, 01-0CT-2000 (TrEMBLrel. 15, 10-UTN-2003 (TrEMBLrel. 24, 3) Surface protein PspC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 58.59
Matches 175; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                          SEQUENCE
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121 STKPEASDTAKPNKPTEPGEKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLELEIA 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        181 ESDVEVKKABLELVKVKANEPRDEQKIKQABAEVESKQABATRLKKIKTDREBABBBAKR 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       240
                                                                                                                                                                                              EIAESDVEVKKAELELVKVKANEPRDEQKIKQAEAEVESKQAEATRLKKIKTDREEAEEE 237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 TITVALVNELANJIKANEYLAKIVESTSESQLQILAMMESRSKVDEAVSKFEKDSSSSSSSS 120
106 NFALNLKLSRIKTEYLYGLSVLKEKSEAELPSKAELPSKIKEKLTAAFEHFKKOT---- 160
                                                                                                                           39 EKEUTIQVPISSNRANKSQ-----TEHMRAAKQVDEYIKKKL-----QLDRRKH 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 ENEGATOVPISSNRANESQAEQGEQPKKLDSERDKARKEVEEYVKKIVGESYAKSIKKRH 60
                                                                                                                                                                                                                                                                                                                              238 AKRRADAK------EQGKPKGRAKRGVPGELATPDKKENDAKSSDSSVGEETL 284
                                                                                                                                                                                                                                                                                                                                                                                      263 AKRRAEAKLKEAVEKNTATSEQGKPKGRAKRGALGEPATPDKKENDAKSSDSSVGEETL 321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               203 BIABSDVBVKKABLELVKBEANKPRNEQKVKQAKAKVBSKQABATRLEKIKTDREBABEB
                                                               118 SDSSTKPEASDTAKPNKPTEPGEKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RAEAKLKEAVEKNTATSEQGKPPKPRGKRGALGEQATPDKKENDAKSSDSSVGEETL 296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          50; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=SFEES;

I Tannelli F., Oggioni M.R., Pozzi G.;

I Tannelli F., Oggioni M.R., Pozzi G.;

I Tannelli F., Oggioni M.R., Pozzi G.;

I "Allello Variation in the Highly Polymorphic Locus pspC of Streptococcus pneumoniae.";

I Submitted (MAY-1999) to the EMBL/GenBank/DDBJ databases.

E MBLG AFF154040; AAF73812.1; -...

R GG: 0016020; C:membrane; IEA.

R InterPro; IPR005479; CW_binding.

R InterPro; IPR007479; CW_binding.

R InterPro; IPR00777; GPOS YSIRK.

R Pfam; PF01473; CW_binding_1; 7.

R Pfam; PF01473; CW_binding_1; 7.

R Pfam; PF05650; XSIRK signal; 1.

R TIGRFAMS; TIGR0168; YSIRK signal; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 681;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Streptococcus pneumoniae.
Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
54.6%; Score 763; DB 2;
Best Local Similarity 57.4%; Pred. No. 9.3e-30;
Matches 170; Conservative 35; Mismatches 41;
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2 EPGEKVABAKKKVEBAEKKAKDQKEBDRRNYPTITYKTLELBIAESDVEVKKAELBLVKV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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New York
Y: U.S.
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                  RESULT 11
US-08-714-741-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
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                                                                                                                                                                                                                                                                                                                                                              2 BPGEKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLELEIAESDVEVKKAELELVKV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 38, Application US/09536784
Patent No. 6573082
GENERAL INFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 81.5%; Score 433; DB 4; Length 453; l Similarity 82.1%; Pred. No. 5.3e-33; 87; Conservative 15; Mismatches 4; Indels
                                                                                                                                                                                                                                                                    Query Match
81.5%; Score 433; DB 3; Length 453;
Best Local Similarity 82.1%; Pred. No. 5.3e-33;
Matches 87; Conservative 15; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            148 EAKEPRNEEKVKQAKAEVESKKAEATRLEKIKTDRKKAEEBAKRKA 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                 62 KANEPRDEOKIKQAEAEVESKQAEATRLKKIKTDREEAEEBAKRRA 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS Version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-0ct-1997
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Human Genome Sciences, Inc. STREET: 9410 Key West Avenue CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION UNDERS: 06/961,083
FILING DATE: 0CT-30-197
ATTORNEY/AGENT INFORMATION:
NAME: MICHOELLE &: Marks
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
TELECHONE: (301) 309-8504
TELEFAX: (301) 309-8512
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MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 38:
US-09-536-784-38
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LENGTH: 453 amino acids
TYPE: amino acid
STRANDEDNESS: single
TELECOMMUNICATION INFORMATION:
                   TELEPHONE: (301) 309-8504
TELEPAX: (301) 309-8512
INPORTATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 38:
                                                                                                           LENGTH: 453 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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COUNTRY: USA
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Best Local Similarity
Matches 87; Conservat
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APPLICANT: Briles, David E.
APPLICANT: McDaniel, Larry S.
APPLICANT: Swiatlo, Edwin
APPLICANT: Yother, Janet
APPLICANT: Crain, Marilyn J.
APPLICANT: Tart, Rebecca
APPLICANT: Tart, Rebecca
APPLICANT: Tart, Rebecca
APPLICANT: Tart, Rebecca
APPLICANT: Brooks Walter, Alexis
TITLE OF INVENTION: EXPRESSION PRODUCTS THEREPROM, AND USBS OF SUCH GENES,
TITLE OF INVENTION: PORTIONS AND PRODUCTS
88 KPEKKVABABEKKKVEBAKKKABDQKEBDRANYPINTYKTLELEIABSDVBVKKABLELVKE 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 EKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLELEIAESDVEVKKAELELVKVKAN 64
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                                                                                                                62 KANEPRDEOKIKOAEAEVESKOAEATRLKKIKTDREEAEBEKRRA 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                65 EPRDEOKIKOAEAEVESKOAEATRIKKIKTDREEAEEEAKRRADA 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READBRIE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE: US/08/714,741
FILING DATE: 16-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Curtis, Morris & Safford, P.C. 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      454312-2460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Frommer Esq., William S. REGISTRATION NUMBER: 25,506
                                                                                                                                                                                                                                                                                  Sequence 40, Application US/08714741; Patent No. 6500613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 454
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 40;
SEQUENCE CHARACTERISTICS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 12
US-09-286-981B-10
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us-09-056-019b-9.rai

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MOLECULE TYPE: amino acid US-08-714-741-46
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   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New York : U.S.
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                  172 KEKVESKKAEAIRLENIKIDRKKA-EEEAKRKADAKLKEANVAISDQGKPKGRAKRGVPG 230
                                                                                                                                                                                                                                                                                                                                                                        115 EAKKKAKAQKEEDHRNYPTITYKTLDLEIAEFDVKVKEABLELVKKEADESRNEGTINQA 174
                                                                                                                                                                                                    61 KYLRELNVLEEKSKOELPSEIKAKLDAAFEKFKKOTL--KPGEKVAEAKKKVE---- 111
                                                                                                                                                                                                                                                                           112 BAKKKAEDOKEEDRRNYPINTYKTLELEIAEFDVKVKEAELELVKEEAKESRNEGTIKQA 171
                                                                                                                              1 ENEGSTOAATSSNMAKTEHRKAAKOVVDEYIEKMIREIQLDRRKHTQNVALNIKLSAIKT 60
                                                                                                                                                   2 ENBRITQVPTSSNRGKPERRKAAAQ-FDEYINKM---IQLDKRKHTQNLAFNIQLSRIKT 57
                                                                                            Gaps
                                                   Query Match

66.1%; Score 826.5; DB 4; Length 419;
Best Local Similarity 70.1%; Pred. No. 1.1e-57;
Matches 185; Conservative 26; Mismatches 30; Indels 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PNEUMOCOCCAL GENES, PORTIONS THEREOF
EXPRESSION PRODUCTS THEREPROM, AND US
PORTIONS AND PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30
URRENT APPLICATION DATA:
APPLICATION UNMER: US/08/714,741
FILING DATE: 16-SEP-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 : Curtis, Morris & Safford, P.C. 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
REPERNCE/DOCKET NUMBER: 454312-2460
TELECOMMUNICATION INFORMATION:
TELECHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                              229 EQATPDKKENDAKSSDSSVGEETL 252
                                                                                                                                                                                                                                                                                                                                                                                                                           231 ELATPDKKENDAKSSDSSVGEETL 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
; ORGANISM: Streptococcus pneumoniae US-09-286-981B-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Brooks-Walter, Alexis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 46, Application US/08714741
Patent No. 6500613
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Crain, Marilyn J.
Hollingshead, Susan
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Swiatlo, Edwin
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SECUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New York
: New York
RY: U.S.
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APPLICANT: Briles
APPLICANT: McDani
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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US-08-714-741-46
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single

STRANDEDNESS:

LENGTH:

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THEREFROM, AND USES OF SUCH GENES,
                                                                                                                                                                                                                                                                                                                                                                                   226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   287 VKEEAXEXRNEEKIKOAKEKVESKKAEATRIEKIKTDRKKAEEEBAKRKAEESEKKA--AE 344
                                                                                                                                                                 47 AKTEHRKAAKKVVDEYIEKMLREIQLDRRKHTONVALNIKLSAIXTKYLRELSVLKENSK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    163 -----RNEGTIKQAKEKVESKKAEATRLENIKTDRKKAEBEAKRKADAKLKEANVAT 214
                                                                                                                                                                                                                                                                                                                                  85 LDAAFEKFKKUTLKPGEKVABAKKKVEBAKKKAEDOKEEDRRNYPTNTYKTLELEIAEFO 144
                                                                                                                                                                                                                        70 -----EEKSKOELPSEIKAK 84
                                                                                                                                                                                                                                                                          107 KEELTSKTKAELTAAFEQFKKDTLKPEKKVAEAEKKVEEAELXVXEEKSXXELPSEIKAK
                                                                                                                                                                                                                                                                                                                                                                        167 LDAAFXKFKKOTLKPGEKVAEAKKXVEEAKKKAKOQKEEDRRNYPTNTYKTLELBIABSD
                                                                                                               15 AKTEHRKAAKQVVDEYIERMLREIQLDRRKHTQNVALNIKLSAIKTKYLRELNVL----
                                                       28; Indels 106; Gaps
     Length 605;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              215 SDQGKPKGRAKRGVPGELATPDKKEN---DAKSSD 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    345 ADEEKIK-QAKAKVESKKAEATRLENIKTDXKKAE 378
Query Match 64.3%; Score 805; DB 4; Best Local Similarity 56.7%; Pred. No. 8.4e-56; Matches 190; Conservative 11; Mismatches 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3: Curtis, Morris & Safford, P.C.
530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Tart, Rebecca
APPLICANT: Brooks-Walter, Alexis
TILE OF INVENTION: PNEUMOCOCCAL GENES, PC
TILLE OF INVENTION: EXPRESSION PRODUCTS IF
TILLE OF INVENTION: PORTIONS AND PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            454312-2460
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MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5-08-714-741-40
Sequence 40, Application US/08714741
Patent No. 6500613
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NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
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McDaniel, Larry S.
Sviatlo, Edwin
Yother, Janet
Crain, Marilyn J.
Hollingshead, Susan
Tart, Rebecca
Brooks-Walter, Alexie
                                                                                                                                                                                                                                                                                                                                                                                                                                           145 VKVKEAELELVKEEAKES----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 45
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(212) 840-0712
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229 TLPSPSLIMMANESQTEHRKD----VDEYIKKMLSEIQLDRRKHTQNVNLNIKLSAIKTKY 284
                                                                                                                                                                                                                                                                                                                                                                                                              63 LRELNVLEEKS-KDELPSEIKAKLDAAFEKFKKDTLKPGEKVAEAKKKVEBAKKKAEDOK 121
                                                                                                                                                                                                                                                                                                                                                                                                                                             285 LYELSVLKENSKKEELTSKTKAELTAAFEQFKKOTIKPEKKVAEAEKKVEEAKKVEEAKKAKAKOQK 344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          122 BEDRRNYPTNTYKTLELEIAEFDVKVKEAELELVKEEAKESRNEGTIKQAKEKVESKKAE 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           182 ATRIENIKTDRKKAEEEAKRKADAKUKEANVATSDQGKPKGRAKRGVPGELATPDKK--- 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ------AEAKOKVDAEEYALEAKIAE 453
                                                                                                                                                                                                                                                                                                                     6 TOAATSSNWA---KTEHRKAAKOVVDEYIEKMLREIQLDRRKHTQNVALNIKLSAIKTKY
                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                   Query Match 61.6%; Score 770; DB 4; Length 864; Best Local Similarity 67.0%; Pred. No. 7.5e-53; Matches 177; Conservative 18; Mismatches 39; Indels 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          405 ATRLEKIKTDRKKABERAKRKABESEKKA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       239 -----ENDAKSSDSSVGEETL 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | : | | : | 454 LEYEVQRLEKELKEIDESDSEDYL 477
information for SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 864 amino acids
TYPE: amino acid
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: amino acid
US-08-714-741-40
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Search completed: March 8, 2004, 12:37:48 Job time : 11.7102 secs

us-09-056-019b-10.rai

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New York
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         ADDRESSEE:
STREET: 53C
CITY: New Y
STATE: New
COUNTRY: U.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 KSGKKVAEAEKGKVEBAEKGAKDQKEEDRRNYPTNTYKTLDLEIAESDVKVKEAELELVKE 60
                                                       APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ô
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 453;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Indels
                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33 OPERAINO SYSTEM: MSDOS Version 6.2 SOFTWARE: ASCII Text CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 468; DB 4;
Pred. No. 1.7e-35;
8; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/961,083
FILING DATE: 0CT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: MIChelle S. Marke:
RESISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULA TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 38:
US-09-536-784-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
Application US/09536784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 38 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM
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Best Local Similarity 89.6
Matches 95; Conservative
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CAL GENES, PORTIONS THEREOF,
NA PRODUCTS THEREFROM, AND USES OF SUCH GENES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       348 KPEKKVABABKKVEBAKKKAKDOKEEDRRNYPINIYKTLELEIAESDVKVKEAELELVK 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 KSGKKVAEAEKKVEEAEKKAKDQKEEDRRNYPINTYKTLDLEIAESDVKVKEAELELVKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4; Length 588;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 EAKEPRDEEKIKQAKAKVESKKAEATRLENIKTDRKKAEEEAKRKA 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: PERFORM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
APPLICATION DATA:
RILING DATE: 16-SEP-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.8e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 90.3%; Score 467; DE Best Local Similarity 91.5%; Pred. No. 2.8e Matches 97; Conservative 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: PNEUMOCOCCAL GENES, PO
TITLE OF INVENTION: EXPRESSION PRODUCTS TO
TITLE OF INVENTION: PORTIONS AND PRODUCTS
                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fornmer Esq., william S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454312-TELECOMMINICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 40, Application US/08714741
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (212) 840-3333
530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (212) 840-333
TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACECULE TYPE: amino acid US-08-714-741-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            530 Fifth Avenue
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321 KPEKKVABABKKVEBAKKKAKDOKBEDRRNYPINTYKTLELBIABSDVKVKBABLELVKE 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 KSGKKVABABKKVEBABKKAKDQKEBDRRNYPTNTYKTLDLBIABSDVKVKBABLELVKE 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 KKVAEAEKKVEBAEKKAKDQKEEDRRNYPINTYKTLDLEIAESDVKVKBAELELVKEBAK 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 KKVAEAEKKVEEAKKKAEDOKEEDRRNYPTNTYKTLELEIAESDVEVKKAELELVKEEAK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Wizemann, Theresa M.
APPLICANT: Wizemann, Theresa M.
APPLICANT: Koenig, Scott
APPLICANT: Koenig, Scott
APPLICANT: Johnson, Leslie S
TITLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
FILE REFERENCE: 469201-396
CURRENT APPLICATION NUMBER: US/09/286,981B
CURRENT APPLICATION NUMBER: US 60/085,743
PRIOR APPLICATION NUMBER: US 60/085,743
PRIOR PILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 1
LENGTH: 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 89.6%; Score 463; DB 4; Length 103; Best Local Similarity 92.2%; Pred. No. 8.9e-36; Matches 95; Conservative 6; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 90.3%; Score 467; DB 4; Length 864; Best Local Similarity 91.5%; Pred. No. 4.3e-35; Matches 97; Conservative 3; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 BAKEPRDEEKIKQAKAKVESKKABATRLENIKTDRKKAEEBAKRKA 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 381 BANESRNEEKIKQAKEKVESKKAEATRIEKIKTDRKKAEEEAKRKA 426
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-1996
CLASSIFICATION: 45-1996
ATTORNEY/AGENT INFORMATION:
NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 45-4312-2460
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 840-772
INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS:
LENGTH: 68-4 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09286981B Patent No. 6503511
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear

MOLECULE TYPE: amino acid
US-08-714-741-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-286-981B-1
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8, 2004, 12:37:49

completed: March me : 5.46962 secs

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88.4%; Score 464; DB 4; Length 864;
  (212) 840-0712
                                                                                            TYPE: amino acid
STRANDEDNESS: sir
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TELEFAX:
                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Brides, David E.
APPLICANT: Brides, David E.
APPLICANT: McDaniel, Larry S.
APPLICANT: Swiatlo, Edwin
APPLICANT: Yother, Janet
APPLICANT: Crain, Marilyn J.
APPLICANT: Tart, Rebecca
APPLICANT: Brooks-Walter, Alexis
TITLE OF INVENTION: PREPRESSION PRODUCTS
TITLE OF INVENTION: PORTIONS AND PRODUCTS
TITLE OF INVENTION: PORTIONS AND PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100 PGEKVABAKKKVEBAKKKABDQKBEDRRNYPINIYKILELBIAEFDVKVKBAELELVKBE 159
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                                                                                    APPLICANT: Wizemann, Theresa M.
APPLICANT: Koenig, Scott
APPLICANT: Toolnago, Scott
APPLICANT: Toolnago, Scott
TITLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
FILE REFERENCE: 469201-396
CURRENT APPLICATION NUMBER: US/09/286,981B
CURRENT FILING DATE: 1999-04-06
PRIOR APPLICATION NUMBER: US 60/085,743
PRIOR PILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
LENGTH: 251
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-1996
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Curtis, Morris & Safford, P.C. 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-09-286-981B-4
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NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 45431
TELECOMMUNICATION INFORMATION:
                 Sequence 4, Application US/09286981B
Patent No. 6503511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 42, Application US/08714741
Patent No. 6500613
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New York
: New York
RY: U.S.
                                                                GENERAL INFORMATION:
APPLICANT: Wizemann
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US-08-714-741-42
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349 PEKKVABAEKKVEBAKKXAKOQKEEDRRNYPINIYKTLELEIABSDVKVKEABLELVKEE 408
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ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYER: Rloppy disk
COMPUTER: RAPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFFWARE: PACENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-196
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454312-2460
TELEPAX: (212) 840-3133
TELEPAX: (212) 840-0712
INFORMATION POR SEQ ID NO: 40:
SEGURNCE CHRACTERISTICS:
TENETH READAMERISTICS:
TENETH READAMERISTICS:
TENETH READAMERISTICS:
TENETH READAMERISTICS:
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APPLICANT: Yother, Janet
APPLICANT: Crain, Marilyn J.
APPLICANT: Crain, Marilyn J.
APPLICANT: Tark, Rebecad
APPLICANT: Tark, Rebecad
APPLICANT: Brooks-Walter, Alexis
TITLE OF INVENTION: EXPRESSION PRODUCTS
TITLE OF INVENTION: EXPRESSION PRODUCTS
TITLE OF INVENTION: PORTIONS AND PRODUCTS
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris & Safford, P.C.
STREET: 530 Fifth Avenue
CITY: New York
STATE: New York
CUNTRY: U.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40, Application US/08714741 Patent No. 6500613 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Briles, David E. APPLICANT: McDaniel, Larry S. APPLICANT: Swiatlo, Edwin
                                                                                                                                           TOPOLOGY: linear

MOLECULE TYPE: amino acid
US-08-714-741-42
                                                          588 amino acids
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                        322 PEKKVABABAKKVERAKKKAKDQKEEDRRNYPTNTYKTLELRIASSDVKVKBABLELVKES 381
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wizemann, Theresa M.
APPLICANT: Wizemann, Theresa M.
APPLICANT: Kooning, Scott
APPLICANT: Kooning, Scott
TILLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
TILLE PERBERGER, 469210-396
CURRENT APPLICATION NUMBER: US/09/286,981B
CURRENT APPLICATION NUMBER: US/09/286,981B
CURRENT PELLING DATE: 1999-04-06
FRIOR APPLICATION NUMBER: US/09/286,981B
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 15
LENGTH: 419
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                                                                              1 PGEKVABAKKKVEBAKKKABDQKEBDRRNYPINTYKTLELBIABFDVKVKBAELELVKEB 60
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APPLICANT: Wizemann, Theresa M.
APPLICANT: Koenig, Scott
APPLICANT: Conig, Scott
TITLE OF INVENTION: Derivatives of Choline Binding Froteins for Vaccines
FILER REPRENCE: 469201-396
CURRENT APPLICATION NUMBER: US/09/26,981B
CURRENT FILING DATE: 1999-04-06
PRIOR FILING DATE: 1999-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
LENGTH: 103
                             0; Gaps
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86.1%; Score 452; DB 4; Length 103;
Best Local Similarity 91.3%; Pred. No. 1.5e-34;
Matches 94; Conservative 4; Mismatches 5; Indels
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Best Local Similarity 90.6%; Pred. No. 1.4e-34;
Matches 96; Conservative 4; Mismatches 6; Indels
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Sequence 15, Application US/09286981B

Setter No. 6503511

GENERAL INFORMATION:
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Patent No. 6503511
GENERAL INFORMATION:
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APPLICANT: Wizemann, Theresa M.
APPLICANT: Wizemann, Theresa M.
APPLICANT: Wordson, Scott
APPLICANT: Johnson, Leslie S
FILE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
FILE REFERENCE: 469201-396
FILE REFERENCE: 469201-396
FILE REFERENCE: 1999-04-06
FILE REFERENCE: 1999-04-06
FILE REFERENCE: 1999-04-06
FILE REFERENCE: 1999-05-16
FILE REFERENCE: 1999-05-18
FILE REFERENCE: 1999-05-18
FILE REFERENCE: PROTEIN NUMBER: US 60/085,743
FRICE FILE REFERENCE: 1998-05-18
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
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                                   APPLICANT: Wisemann, Theresa M.
APPLICANT: Wisemann, Theresa M.
APPLICANT: Koenig, Scott
APPLICANT: Condig, Scott
APPLICANT: Condig, Scott
CURRENT: Johnson, Leelie S
FILE REFERENCE: 469201-396
FILE REFERENCE: 469201-396
FILE REFERENCE: 1999-04-06
PRICE APPLICATION NUMBER: US 60/085,743
PRICE APPLICATION NUMBER: US 60/085,743
PRICE RILNG DATE: 1999-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
3 EKVABAKKKVEBAKKKABDOKEBDRRNYPINIYKTIBIBIABFDVKVKBABIBIVKBBAK 62
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85.5%; Score 449; DB 4; Length 428;
Best Local Similarity 87.6%; Pred. No. 1.46-33;
Matches 92; Conservative 7; Mismatches 6; Indels
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86.1%; Score 452; DB 4; Length 114;
Best Local Similarity 91.3%; Pred. No. 1.7e-34;
Matches 94; Conservative 4; Mismatches 5; Indels
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                                                                                                                            63 ESRNEGTIKQAKEKVESKKAEATRLENIKTDRKKAEEEAKRKA 105
                                                                                                                                                                       61 ESRNEEKIKQAKAKVESKKAEAIRLEKIKTDRKKAEEEAKRKA 103
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                                                                                                                                                                                                                                                                                                                                   Sequence 19, Application US/09286981B Patent No. 6503511 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/09286981B; Patent No. 6503511; GENERAL INFORMATION:
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ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                     RESULT 10
US-09-286-981B-19
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US-09-286-981B-16
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APPII
                                                           March 8, 2004, 12:22:33 ; Search time 5.14427 Seconds (without alignments) 1224.345 Million cell updates/sec
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                                                                                                          593
1. PSSSLKSGKKVAEAEKKVEE......AEEEAKRKAAEEDKVKEKRA 122
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Sequence 3
Sequence 5
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Sequence
Sequence
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/cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
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/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
        GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-08-84-1065-25
US-09-308-025-6
US-09-286-981B-10
US-09-286-981B-12
US-09-286-981B-13
US-09-286-981B-7
US-09-286-981B-7
US-09-286-981B-7
US-09-286-981B-7
US-09-286-981B-10
US-09-286-981B-11
US-09-286-981B-11
US-09-286-981B-11
US-09-286-981B-11
US-09-286-981B-13
                                                                                                                                                                                 Total number of hits satisfying chosen parameters:
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US-09-286-981B-38
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                                                                                                                                                                  389414 segs, 51625971 residues
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Maximum Match 100%
Listing first 45 summaries
                                          OM protein - protein search, using sw model
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Match Length
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Sequence 16, Application US/09286981B
Patent No. 6503511
Fatent No. 6503511
Fatent No. 6503511
APPLICANT: Wizemann.Theresa M.
APPLICANT: Wizemann.Theresa M.
APPLICANT: Kenigh Scott
FILE REFERENCE: 469201-36
FILE REFERENCE: 469201-36
FILE REFERENCE: 469201-36
FILE REFERENCE: 1999-04-06
FILE REFERENCE: 1999-04-06
FRIOR APPLICATION NUMBER: US 60/085,743
FRIOR APPLICATION NUMBER: US 60/085,743
FRIOR FILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: RatentIn Ver. 2.1
SEQ ID NO 16
IENGTH: 414
TYPE: PRI
CREATE FILE REPROCECTORS pneumoniae
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102, App
57, Appl
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2,9581, A
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6321, A.
Appli
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Patent No. 6245335
GENERAL INFORMATION:
APPLICANT: ABBLICANT: ROBEROW GENERAL INFORMATION:
APPLICANT: TRONGER Eliane
APPLICANT: TRONGER Eliane
APPLICANT: TRONGER Eliane
APPLICANT: TRONGER ELIANE
TITLE OF INVENTION: ANTI-PNEUMOCOCCAL VACCINES
US-08-714-741-41

US-08-529-055-56

US-08-714-741-32

US-08-714-741-32

US-09-516-784-102

US-09-529-055-57

US-09-328-328-321

US-09-329-329-5

US-09-543-681A-6436

US-09-543-681A-6436

US-09-543-681A-6436

US-09-543-681A-6436

US-09-543-681A-6436

US-09-525-991A-29581

US-08-46-746-2

US-08-214-164-2

US-08-214-164-2

US-08-214-164-2

US-08-214-164-2

US-08-214-164-2
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Pred. No. 6.2e-46;
0; Mismatches 1;
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Best Local Similarity 99.2
Matches 121; Conservative
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